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Result
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Listing first 45 summaries
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1: /SIDS1/gcgdata/hold-geneseq/geneseqp-emb1/AA1980.DAT:*

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3: /SIDS1/gcgdata/hold-geneseq/geneseqp-emb1/AA1983.DAT:*

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(first entry)

Screening for ligand analogs and agents which modulate ligand-receptor binding, comprises adding a test ligand to a non-naturally occurring cell surface receptor analog - $\,$ 11-FEB-1999; 29-APR-1999; 11-FEB-2000; 2000WO-US03665 WO200047612-A2. Homo sapiens. Ligand; cell surface receptor; erythropoietin; EPOR; human; protein design automation; PDA. Human mature Luo P, (XENC-) XENCOR INC 17-AUG-2000 2000-549135/50 Dahiyat B; erythropoietin receptor EPOR extracellular domain 99US-0120009 99US-0131674 #2

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Matches 198
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present invention relates to a method for screening for a ligand log, comprising adding a candidate ligand to a non-naturally occurring surface receptor analog e.g. erythropoietin receptor (EPOR), and
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                                                                                        which modulate ligand-receptor to a non-naturally occurring
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Best Local
      The present sequence is truncated human EpoR (erythropoietin receptor). Transfection of constitutively active EpoR(e439) by electroporation into a cytokine-dependent cell line supports cell population expansion in the absence of exogenous cytokines. Mutant human EpoR is used in treatment of disorders related to inadequate EpoR signalling. The transfected cells
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                                                                                                                                                                                                                                                                                                                                          Truncated human EpoR; erythropoietin receptor; hypersensitive EpoR(t439); mutant human EpoR; EpoR signalling; cancer; infectious disease; HIV; sickle cell anaemia; cytostatic; antimicrobial; antiviral;
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25-JAN-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            225;
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                                                                                                           stem
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also

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to

treat cancer,

infectious

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AAWOB3449
ID AAWO
XX AAWO
XX AAWO
XX EpoR
XX EpoR
XX Rece
KW immu
DE Chim
OS Chim
OS Chim
OS Chim
YY Key
FT Doma
FT Clea
FT Clea
FT Doma
FT OT-J
XX WO96
XX WPI;
DR N-PS
XX Reco
PT For
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Qy
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Best Local Similarity
Matches 198; Conserv
                                                                                                      07-JUN-1995;
                                                                                                                        07-JUN-1996;
                                                                                                                                                                                                  Domain
                                                                                                                                                                                                                  Cleavage-site
                                                                                                                                                                                                                                                       Domain
                                                                                                                                                                                                                                                                                   Chimeric
                                                                                                                                                                                                                                                                                           Chimeric
                                                                                                                                                                                                                                                                                                              anaemia;
                                                                                                                                                                                                                                                                                                                                                   EpoRFc fusion
                                                                                                                                                                                                                                                                                                                                                                      14-MAR-1997
                                                                                                                                                                                                                                                                                                                                                                                                         AAW08349 standard;
Recombinant immunogen corresp. to dimeric form of a receptor - used for generating antibodies able to act as receptor agonists, esp. of erythropoietin receptor for treating anaemia
                                                                  Erickson-Miller
                                                                                                                                           19-DEC-1996
                                                                                                                                                            W09640231-A1
                                                                                                                                                                                                                                                                                                                      Receptor agonist; antibody; erythropoietin receptor; immunogen; antigen; metallothionein; promoter; IgG1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          expression
                                                                                   (SMIK ) SMITHKLINE BEECHAM CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                                                        214
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                                               1997-051900/05
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            HQAPTARGAIRFWCSLPTADTSSFVPLELRLTAASGAPRFHRVIHINEVVLLDAPVGLVA 120
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TRITIAVRARMAEPSFGGFWSAWSEPVSLLT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RLADESGHVVIRWLPPPETPMTSHIRFELDISAGNGAGSVQRVELLEGRTECVLSNLRGR 180
                                                                                                                                                                                                                                                                                                                                                                                                                                                       trytfavrarmaepsfggfwsawsepvsllt
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         {\tt rladesghvvlrwlpppetpmtshiryevdvsagngagsvqrveilegrtecvlsnlrgr}
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         HIV), sickle cell anaemia, ssion of erythropoietin.
                                       AAT48800.
                                                                                                                                                                                                                                                                                 synthetic.
                                                                                                                                                                                                                                                                                                            therapy.
                                                                                                                                                                                                                                                                                           Homo sapiens;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        438
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                    (first entry)
                                                                                                                                                                                                                                                                                                                                                   protein
                                                                CL,
                                                                                                     95US-0474673.
                                                                                                                        96WO-US09613
                                                                                                                                                                                                /note= "E
255..488
                                                                                                                                                                                                                  251..254
                                                                                                                                                                                                                          Location/Qualifiers
                                                                                                                                                                              /note=
                                                                                                                                                                                         ′label=
                                                                                                                                                                                                                                                                                                                                                                                                         Protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 82.6%; Score 1060; 93.8%; Pred. No. 2
                                                                                                                                                                              "human
                                                                                                                                                                                                         "Factor Xa
                                                                                                                                                                                         FC
                                                                                                                                                                                                                                                                                                                                                                                                         488
                                                                                                                                                                               IgG1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                         B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   and conditions related to abnormal
                                                                                                                                                                                                        cleavage site'
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DB 21;
2.1e-98;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length
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RESULT
AAB13012
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A fusion protein (AAW08349) encoded by plasmid mtalsEpoRFc (AAT48800) comprises the human erythropoietin receptor (EpoR) extracellular domain fused (via a Factor Xa cleavage sequence) to the Fc portion of human IgG1. It can be expressed e.g. in transfected brosophila $2 cells upon induction with copper sulphate. The cells secrete EpoRFc as a dimeric molecule due to the affinity of the Fc moiety for itself. The dimeric receptor can be used as an immunogen to generate antibodies (monoclonal, polyclonal, chimeric, humanised) able to act as EcoR agonists for use in treatment of anaemia.
                                                                                WPI;
                                                                                                     Tew
                                                                                                                                                                                                          WO200043492-A2.
                                                                                                                                                                                                                                                 erythropoietin
                                                                                                                                                                                                                                                           Site specific label; detection;
                                                                                                                                                                                                                                                                                Q-tagged erythropoietin (EPO) receptor protein.
                                                                                                                                                                                                                                                                                                     08-DEC-2000
                                                                                                                                                                                                                                                                                                                         AAB13012;
                                                                                                                                                                                                                                                                                                                                             AAB13012 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence
                                                                                                                                             22-JAN-1999;
                                                                                                                                                                 20-JAN-2000;
                                                                                                                                                                                     27-JUL-2000
                                                                                                                                                                                                                              Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 1; Page 39-41; 83pp;
                                                                                                                        (SMIK )
                                                                                                                                                                                                                                                                                                                                                                                                 214
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                                                                                2000-499222/44
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                                                                                                                                                                                                                                                                                                                                                                                                trytfavrarmaepsfggfwsawsepvsllt
                                                                                                                                                                                                                                                                                                                                                                                                                      TRITIAVRARMAEPSFGGFWSAWSEPVSLLT 211
                                                                                                                                                                                                                                                                                                                                                                                                                                       \verb|rladesghvvlrwlpppetpmtshiryevdvsagngagsvqrveilegrtecvlsnlrgr|
                                                                                                                                                                                                                                                                                                                                                                                                                                                  RLADESGHVVIRWLPPPETPMTSHIRFELDISAGNGAGSVQRVELLEGRTECVLSNLRGR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  198;
                                                                                                                        SMITHKLINE BEECHAM CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Similarity
                                                                                                    Powell DJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       488 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                  2000WO-US01481
                                                                                                                                                                                                                                                                                                     (first entry)
                                                                                                                                              99US-0117327
                                                                                                                                                                                                                                                 receptor;
                                                                                                                                                                                                                                                                                                                                              Protein;
                                                                                                     Meek TD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            82.6%; Score 1060; 93.8%; Pred. No. 2
                                                                                                                                                                                                                                                  EPO.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  11;
                                                                                                                                                                                                                                                                                                                                              503
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                                                                                                     Chen
                                                                                                                                                                                                                                                           interaction
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                                                                                                     Σ
                                                                                                                                                                                                                                                                                                                                                                                                 244
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DB 18;
.5e-98;
                                                                                                                                                                                                                                                           screening; transglutaminase;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      488;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                             180
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           93
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              60
                                                                                                                                                                                                                                                                                                                                                                                                                                           213
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0;
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This invention relates to methods for

the site specific modification

of

Example 4; Page

26; 49pp;

English

Screening for a candidate compound for use in bioassays comprises contacting the candidate molecule with a labelled modified protein detecting the label to identify interaction of the two molecules -

and

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CCCCCXXXTT PTIXX
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            RESULT
AAR06512
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local S
Matches 198
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           a protein, and to a method for screening for a candidate compound which interacts with first protein. The screening method comprises contacting the candidate molecule with a labelled modified first protein and detecting the label to identify interaction of the labelled modified first protein and candidate compound. The first protein is modified to contain a peptide, represented by sequence AAB13005. The method is used to label proteins at specific sites. The present sequence represents a Q-tayged erythropoletin (EPO) receptor constructed in an example of the method of the invention.
The sequence was deduced from DNA obtd. from a clone isolated a commercially available human geenomic cDNA library in phage Lambda Fix (Stratagene). The sequence encodes a type I transmembrane protein with binding affinity for EPO. The gene and
                                                                                                                                                                                                                                                                                                                                                       D'andrea
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       01-FEB-1990;
                                                                                                                                   Disclosure; Fig
                                                                                                                                                                                        and systems
                                                                                                                                                                                                                Erythropoietin receptor and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        03-FEB-1989;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              09-AUG-1990
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WO9008822-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Erythropoietin;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         EPO receptor
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAR06512;
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                                                                                                                                                                                                                                                                                                                                                                                                           (WHIT-)
                                                                                                                                                                                                                                                                                                                                                                                                                                    (GENE-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       214
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DB; AAQ05748.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RLADESGHVVIRWLPPPETPMTSHIRFELDISAGNGAGSVQRVELLEGRTECVLSNLRGR
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     rladesghvvlrwlpppetpmtshiryevdvsagngagsvqrveilegrtecvlsnlrgr
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                  Wong
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          90WO-US00635
                                                                                                                                   2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Diamond
                                                                                                                                                                                                                                                                                                                                                    G
                                                                                                                                 53pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           82.6%; Score 1060; DB 21;
93.8%; Pred. No. 2.6e-98;
tive 11; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Blackfan
                                                                                                                                   English
                                                                                                                                                                                                                gene
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        polycythemia
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Best Local s
Matches 198
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           recombinant EPO receptor produced on expression of the DNA are used to develop reagents and systems to control and study erythropoiesis. It is believed that the EPO receptor is dysfunctional in individuals with Diamond Blackfan anaemia, and may be hyperactive in polycythemia vera. See also AAR06511 (murine EPO receptor).
cleavage site
        expression as a
              New pure
                               N-PSDB;
                                                     Lee
                                                                                                   15-AUG-1994;
                                                                                                                                  WO9505469-A
                                                                                                                                                         Misc-difference
                                                                                                                                                                         Misc-difference
                                                                                                                                                                                        Domain
                                                                                                                                                                                                       Domain
                                                                                                                                                                                                               Key
                                                                                                                                                                                                                                            Erythropoietin
                                                                                                                                                                                                                                                           Human erythropoietin receptor
                                                                                                                                                                                                                                                                           07-OCT-1995
                                                                                                                                                                                                                                                                                          AAR70032;
                                                                                                                                                                                                                                                                                                          AAR70032
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence
                                                                    (LEEJ/)
                                                                                    16-AUG-1993;
                                                                                                                    23-FEB-1995
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                                                                                                                                                                                                                                                                                                                                                                            il Similarity
198; Conserv
                                                                     EEE
                              AAQ82990
               human erythropoietin receptor fragment -
                                                                                                                                                                                                                                                                                                         standard;
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                                                                                                                                                                                                                                                                           (first entry)
                                                                                                                                                                                                                                            receptor; extracellular domain.
                                                                                     93US-0106815
                                                                                                   94WO-US09298
        fusion
                                                                                                                                                        /note= "forward 222..226
                                                                                                                                                                      /note= "extracellular 25...29
                                                                                                                                                                                        /note=
9..83
                                                                                                                                                                                                      Location/Qualifiers 25..250
                                                                                                                                                  /note=
                                                                                                                                                                                                                                                                                                          Protein;
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93.8%;
        protein
                                                                                                                                                                                                "extracellular
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 1060; I
Pred. No. 2.66
11; Mismatches
                                                                                                                                                                                                                                                                                                          508
        having a thrombin
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                                                                                                                                                                 primer
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                                                                                                                                                                                                                                                                                                                                                              211
                                                                                                                                                                                domain"
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                                                                                                                                                  AAQ82992 specific
                                                                                                                                                                 AAQ82991 specific site"
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The full-length erythropoietin receptor (EPO-R) is given. Extracellular domains are expressed from vector plasmid pGEX-2T as fusion proteins with glutathione-S-transferase. The domains are used for investigating the structure of the EPO-R and for identifying factors involved in regulating differentiation and proliferation mechanisms in erythroid progenitor cells. They can also be used for identifying and quantitating EPO and EPO-R as well as in understanding haematopoietic malignancy and some
                                                                                            03-FEB-1989;
25-MAR-1991;
                                                                                                                                                                03-FEB-1989;
                                                                                                                                                                                                              03-JAN-1995
                                                                                                                                                                                                                                                          US5378808-A
                                                                                                                                                                                                                                                                                                                                   Domain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Homo
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human erythropoietin receptor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  11-AUG-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAR69503 standard;
                        (GEMY ) GENETICS
                                                                                                                                                                                                                                                                                                                                                                              Modified
                                                                                                                                                                                                                                                                                                                                                                                                                          Protein
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                                                                      10-JUN-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Peptide
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RLADESGHVVIRWLPPPETPMTSHIRFELDISAGNGAGSVQRVELLEGRTECVLSNLRGR 180
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                                                                 89US-0306503.
91US-0678877.
93US-0075069.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   receptor; anemia therapy; diagnostic.
                                                                                                                                                                89US-0306503
                                                                                                                                                                                                                                                                                                                              /note=
251..2
                                                                                                                                                                                                                                                                                                                                                                            /note=
76..79
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Location/Qualifiers
1..24
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Protein;
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93.8%;
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                                                                                                                                                                                                                                                                                                                                                 "N-glycosylation site"
                                                                                                                                                                                                                                                                                                                                                                                               "mature protein"
                                                                                                                                                                                                                                                                                                        "transmembrane region"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        42pp;
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Pred. No. 2.6e-98;
""smatches 2;
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RESULT
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Matches 198; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The sequence is that of a 55-kDa human erythropoietin receptor. The receptor polypeptide may be used in purification and detection of erythropoietin, and in production of antibodies for anemia therapy. The polypeptide may also be used for treating individuals who are hypersensitive to erythropoietin or who have elevated erythropoietin levels. They may be used in therapy of e.g. primary
  03-FEB-1989;
25-MAR-1991;
                                                                  03-FEB-1989;
                                                                                                                                                     US5278065-A
                                                                                                                                                                                                                 Region
                                                                                                                                                                                                                                                             Protein
                                                                                                                                                                                                                                                                                                                                                                        Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                Erythropoietin
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     24-JUN-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAR47518;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAR47518 standard;
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                                                                                                                                                                                                                                                                                                      Peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence
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91US-0678877
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                                                                  89US-0306503
                                                                                                                                                                                                            /note=
251..2
                                                                                                                                                                                                                                                             /note= "signal 25..508
                                                                                                                                                                                                                                                                                                                          Location/Qualifiers
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93.8%;
                                                                                                                                                                                                                                       "mature
                                                                                                                                                                                             "putative transmembrane
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                                                                                                                                                                                                                                                                                                                                                                                                                   recombinant;
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Pred. No. 2.6e-98;
1; Mismatches 2
                                                                                                                                                                                                                                       EPO receptor"
                                                                                                                                                                                                                                                                                 peptide"
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                                                                                                                                                                                                                                                                                                                                                                                                                   murine;
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RESULT 10
AAY44623
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XX AAY446
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XX TYUNCA
DT 07-APR
DT 07-APR
DT TYUNCA
KW TYUNCA
KW MUTANT
KW Sickle
KW Immuno
XX HOMO S.
XX Key
FH Key
FT Misc-d
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Best Local
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                                                                      Truncated human EpoR; erythropoietin receptor; hypersensitive EpoR(t439); mutant human EpoR; EpoR signalling; cancer; infectious disease; HIV; sickle cell anaemia; cytostatic; antimicrobial; antiviral; immunostimulant; anti-anaemic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Nouse erythroleukaemia (MEL) cells were used to construct a cDNA library. The cDNA was used to transfect COS-1 cells and these were screened for radioiodinated erythropoiethin (EPO) binding to isolate cDNA encoding the EPO receptor. This cDNA was used as a probe to screen a human genomic cDNA library to obtain DNA encoding the human EPO receptor. The cDNA may be used to study, treat or diagnose disorders in which the EPO receptor is dysfunctional The EPO
                                                                                                                                                          07-APR-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       receptor may also be used to raise antibodied or for treating hypersensitivity to EPO or who have elevated levels of EPO. The pord. is pref. used for treating anaemias, primary proliferative polycythemia and secondary polycythemia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Disclosure; Fig
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Recombinant DNA encoding erythropoietin receptor - udevelop prods. for study, treatment or diagnosis of
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              Misc-difference
                                                                                                                                                                                  AAY44623;
                                                                                                                                                                                                       AAY44623 standard; Protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence
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                                                                                                                                   truncated
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GENETICS INST INC.
WHITEHEAD INST BIOMEDIC
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            Location/Qualifiers
                                                                                                                                human EpoR(t439).
 /note=
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92.9%;
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"Wild type Arg substituted by
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                                                                                                                                                                                                                                                                                                                                                                                                                                                        12;
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Pred. No. 1.3e
12; Mismatches
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disorders
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Best Local Similarity 93.4
Conservative
                                                                                                                                            Murine; s
antigen;
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                                                                                                 Mus musculus
                                                                                                                                                                                                                     Mouse soluble
                                                                                                                                                                                                                                                                       19-OCT-1994
                                                                                                                                                                                                                                                                                                                                                                    AAR50326
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence
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25-JAN-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 181 TRITIAVRARMAEPSFGGFWSAWSEPVSLLT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              used
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                                                                                                                                         soluble; erythropoietin; EPO; receptor protein; diagnostic agent; biochemical reagent.
                                                                                                                                                                                                                                                                                                                                                                      standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Matthews KE,
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                                                                                                                                                                                                                                                                       (first entry)
                                                                                                                                                                                                                     EPO receptor protein fragment.
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99CA-2260332.
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                                               Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                      Protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      81.9%;
93.4%;
  "Signal peptide"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mueller
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Pred. No. 1.
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1.4e-97;
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Best Local :
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                                                                                                                          EPO receptor
                                                                                                                                                                                                                                                                                                                                                                                                                                                               This sequence represents a fragment of the murine soluble erythropoietin (EPO) receptor protein (SEPO-R). This protein is able to bind to EPO and has antigenicity as an EPO receptor. The molecular wieght of the full length protein is pref 33 or 29 kD. The protein is useful as a drug, as a diagnostic agent and a biochemical reagent.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI;
          Domain
                                       Domain
                                                        Peptide
                                                                                     Mus musculus
                                                                                                      Erythropoietin; Diamond Blackfan anaemia; polycythemia
                                                                                                                                             04-JAN-1991
                                                                                                                                                                   AAR06511;
                                                                                                                                                                                   AAR06511 standard; protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Soluble erythropoietin receptor protein - SEPO-R, useful as diagnostic reagent
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            04-MAR-1992;
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                                                                                                                                                                                                                                                                                                                                                                                                                                               265
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                                                                                                                                             (first entry)
                                                                                                                          sequence
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                /label=extracellular domain
/note=EPO binding region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      26..265
/note= "Mature EPO-R fragment"
                                              /label=signal peptide
                                                                Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                       68.2%; Score 875.5; DB 15; 77.1%; Pred. No. 4.6e-80;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         9pp; Japanese
                                                                                                                            deduced
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DNA coding
                                                                                                                                                                                                                                                                                                                                                                                              Indels
                                                                                                                                                                                                                                                                                                                                                                                                                 Length
                                                                                                                                                                                                                                                                                                                                                                                                                  265;
                                                                                                                                                                                                                                                                                                                                                                                             1;
                                                                                                                                                                                                                                                                                                                                                                                             Gaps
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                                                                                                                                                                                                                                                                           212
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RESULT 13
AAR47517
ID AAR475
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AC AAR475
XX
DT 24-JUN
XX
XX
DT 24-JUN
XX
XX
XX
EXPENDED MEL EP
XX
KW Erythr

MEL EPO receptor 24-JUN-1994

(first entry)

AAR47517; AAR47517

standard;

Protein;

507

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Erythropoietin receptor; recombinant; murine; anaemia

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                                                                                                                                                                                                                                                                               Matches
                                                                                                                                                                                                                           Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                            dysfunctional in individuals with Diamond Blackfan anaemia, and may be hyperactive in polycythemia vera.

See also AAR06512 (human EPO receptor).
                                                                                                                                                                                                                                                                                                                                                       CDNA library prepd. from uninduced murine crythroleukemia cells. It is a type I transmembrane protein with binding affinity for EPO. The gene and recombinant EPO receptor produced on expression of the DNA are used to develop reagents and systems to control and
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Erythropoietin receptor and gene - used for developing reagents and systems to control and study erythropoiesis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            D'andrea A, Wong G;
                                                                                                                                                                                                                                                                                                                                                                                                             The sequence was deduced from DNA from a clone isolated from
                                                                                                                                                                                                                                                                                                                                                                                                                                       Disclosure; Fig 1; 53pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 1990-260931/34.
N-PSDB; AAQ05747.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              03-FEB-1989;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       01-FEB-1990;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  09-AUG-1990
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (GENE-) GENETICS INST INC (WHIT-) WHITEHEAD INST.
213
            181 TRITIAVRARMAEPSFGGFWSAWSEPVSLLT 211
                                                     153
                                                                 121 RLADESGHVVIRWLPPPETPMTSHIRFELDISAGNGAGSVQRVELLEGRTECVLSNLRGR 180
                                                                                                        93
                                                                                                                                  61
                                                                                                                                                         34 kfeskaallasrgseellcftqrledlvcfweeaassgm-dfnysfsyglegesrkscsl 92
                                                                                                                                                                       KFESKAALLAARGPEELLCFTERLEDLVCFFEEDASAGVGPGNFSFSFQLEDEPWKLCRL
trytfavrarmaepsfsgfwsawsepasllt 243
                                                    rraeegshvvlrwlpppgapmtthiryevdvsagnraggtqrvevlegrtecvlsnlrgg
                                                                                                                 HOAPTARGAIREWCSLPTADTSSFVPLELRLTAASGAPREHRVIHINEVVLLDAPVGLVA 120
                                                                                                    hqaptvrgsvrfwcslptadtssfvplelqvteasgspryhriihinevvlldapaglla
                                                                                                                                                                                                              164;
                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              89US-0306503
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       90WO-US00635
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /label=N-linked_glycos
182..184
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /label=N-linked_glycos
                                                                                                                                                                                                                         67.78;
77.78;
                                                                                                                                                                                                             22;
                                                                                                                                                                                                                           Score 869.5; DB 1
Pred. No. 4.4e-79;
                                                                                                                                                                                                              Mismatches
                                                                                                                                                                                                                                       DB 11;
                                                                                                                                                                                                              24;
                                                                                                                                                                                                              Indels
                                                                                                                                                                                                                                      Length
                                                                                                                                                                                                             1;
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                                                                                                                                                                                   60
                                                                                                        152
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Query Match
Best Local (
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Peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                        Mouse erythroleukaemia (MEL) cells were used to construct a cDNA library. The cDNA was used to transfect COS-1 cells and these were screened for radioiodinated erythropoietin (EPD) binding to isolate cDNA encoding the EPO receptor. The cDNA may be used to isolate the EPO receptor from other sources and to study, treat or diagnose disorders in which the EPO receptor is dysfunctional. The EPO receptor may also be used to raise antibodied or for treating hypersensitivity to EPO or who have elevated levels of EPO. The por is pref. used for treating anaemias, primary proliferative polycythe
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 D'andrea A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      03-FEB-1989;
25-MAR-1991;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           03-FEB-1989;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            11-JAN-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Modified-site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mus musculus.
                                                                                                                                                                                                                                                                                                                                                                          Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Disclosure; Fig 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Recombinant DNA encoding erythropoietin receptor - used to develop prods. for study, treatment or diagnosis of disorders
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 1994-025409/03.
N-PSDB; AAQ53994.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (CHIL-) CHILDRENS MEDICAL CENT. (GEMY) GENETICS INST INC. (WHED) WHITEHEAD INST BIOMEDIC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Modified-site
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213
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       which receptor is dysfunctional
                                    181
                                                                       153
                                                                                                          121
                                                                                                                                            93
                                                                                                                                                                             61
                                                                                                                                                                                                                34
                                                                                                                                                                                                                                                                                                      Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                           secondary polycythemia. also AAR47518.
                                                                                                                                                                                                           RLADESGHVVIRWLPPPETPMTSHIRFELDISAGNGAGSVQRVELLEGRTECVLSNLRGR 180
                                                                                                                                       TRITIAVRARMAEPSFGGFWSAWSEPVSLLT 211
                                                                 rraeegshvvlrwlpppgapmtthiryevdvsagnraggtqrvevlegrtecvlsnlrgg
                                                                                                                                                                                                                                                                                                                                                                          507 AA;
                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Jones SS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    89US-0306503.
91US-0678877.
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383
/note=
250..27
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75
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25..507
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /note=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        24pp; English.
                                                                                                                                                                                                                                                                                                      67.7%; Score 869.5; DB 1 77.7%; Pred. No. 4.4e-79;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                "potential N-glycosylatiom site"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   "potential
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       "signal"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    BIOMEDICAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    "mature EPO receptor"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               "putative transmembrane region"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Wong GG;
                                                                                                                                                                                                                                                                                    22;
                                                                                                                                                                                                                                                                                      Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 N-glycosylation
 243
                                                                                                                                                                                                                                                                                                                     DB 15;
                                                                                                                                                                                                                                                                                      24;
                                                                                                                                                                                                                                                                                      Indels
                                                                                                                                                                                                                                                                                                                     Length 507;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   isolate the
                                                                                                                                                                                                                                                                                                                                                                                                                                          . The pord. polycythemia
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RESULT

AAR695022
ID AAR6

AXX AAR6

AXX AAR6

AXX AAR6

AXX AAR6

AXX MOUS

XXX MUS

ET Pept

FT Pool

FT Mod:

FT Mod:

FT Mod:

FT Mod:

FT Mod:

FT Mod:

FT Doma

FT Mod:

FT OT

AXX US5:

AXX
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Best Local S
                                                                                                                                                                                  Matches 164;
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25-MAR-1991;
10-JUN-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Key
Peptide
                                                                                                                                                                                                                                                                                                                                                        The sequence corresponds to a mouse erythropoietin receptor, including putative signal peptide and transmembrane regions, N-linked glycosylation sites. The protein is derived from merythroleukemia cells and may be used in drug design or in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New recombinant erythropoietin receptor polypeptide(s) - used detection, purificn. and therapy and for prodn. of antibodies detection and therapy % \left( 1\right) =\left\{ 1\right\} 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Erythropoietin receptor; anemia therapy; signal peptide; transmembrane region; N-linked glycosylation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      D'andrea A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mus musculus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mouse erythropoietin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAR69502;
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                                                                                                                                                                                                                                                                                           Sequence
                                                                                                                                                                                                                                                                                                                                        pharmaceutical
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (GEMY)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 03-FEB-1989;
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                                             61
    93
                                                                                           34
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KFESKAALLAARGPEELLCFTERLEDLVCFFEEAASAGVGPGNFSFSFQLEDEPWKLCRL 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1995-051310/07.
                                                                                           kfeskaallasrgseellcftqrledlvcfweeaassgm-dfnysfsyqlegesrkscsl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1; Fig 2;
                                                                                                                                                                                                      Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAQ81891.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GENETICS INST
                                                                                                                                                                                                                                                                                             507
                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Jones SS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first
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91US-0678877.
93US-0075069.
                                                                                                                                                                                                                                                                                             ΑA;
                                                                                                                                                                                                                                                                                                                                      compositions for therapy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 89US-0306503
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383..38
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        24pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /note=
75..77
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25..50
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250..271
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Protein;
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                                                                                                                                                                                                      67.78;
77.78;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          "transmembrane region"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          "mature protein"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        "signal peptide"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               "N-linked glycosylation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Wong
                                                                                                                                                                                  22;
                                                                                                                                                                             Score 869.5; DB 1
Pred. No. 4.4e-79;
2; Mismatches 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GG;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ₽
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f
                                                                                                                                                                                                                           DB 16;
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                                                                                                                                                                                  Indels
                                                                                                                                                                                                                             Length 507;
                                                                                                                                                                                                                                                                                                                                                                                      mouse
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RESULT 15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               QΥ
                                                                                                                                                                   Query Match
Best Local Similarity
Matches 163; Conserv
                                                                                                                                                                                                                                            This sequence represents the murine soluble erythropoietin (EPO) receptor protein (sEPO-R). This protein is able to bind to EPO and has antigenicity as an EPO receptor. The molecular wieght of the full length protein is pref 33 or 29 kD. The protein is useful as a drug, as a diagnostic agent and a biochemical reagent.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Key
Modified-site
                                                                                                                                                                                                                         Sequence
                                                                                                                                                                                                                                                                                                                Disclosure; Fig 1; 9pp; Japanese.
                                                                                                                                                                                                                                                                                                                                     Soluble erythropoietin receptor protein - and DNA coding SEPO-R, useful as diagnostic reagent
                                                                                                                                                                                                                                                                                                                                                                      WPI; 1994-094847/12.
N-PSDB; AAQ44854.
                                                                                                                                                                                                                                                                                                                                                                                                       (SNOW ) SNOW BRAND MILK PROD CO LTD
                                                                                                                                                                                                                                                                                                                                                                                                                             04-MAR-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                   04-MAR-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         15-FEB-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                JP06038787-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mus musculus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Murine; soluble; erythropoietin; EPO; receptor protein; sEPO-R; drug; antigen; diagnostic agent; biochemical reagent.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mouse soluble EPO receptor protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           19-OCT-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAR50327 standard; Protein;
            181
                                             121 RLADESGHVVIRWLPPPETPMTSHIRFELDISAGNGAGSVQRVELLEGRTECVLSNLRGR 180
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 213
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             153
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       181
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  121
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                                                                                                                                            \vdash
                                                                                                                    TRITIAVRARMAEPSFGGFWSAWSEPVSLLT 211
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RLADESGHVVIRWLPPPETPMTSHIRFELDISAGNGAGSVQRVELLEGRTECVLSNLRGR 180
                                rraeegshvvlrwlpppgapmtthiryevdvsagnraggtqrvevlegptecvlsnlrgg
                                                                                                                                                                   163; Conservative
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                                                                                                                                                                                                                         507 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                   92JP-0082865
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Location/Qualifiers 75..77
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /note= "N-linked glycosylation site"
                                                                                                                                                                67.2%; Score 862.5; DB 15; Length 507; 77.3%; Pred. No. 2.3e-78; tive 22; Mismatches 25; Indels 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         507
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Db 213 trytfavrarmaepsfsgfwsawsepasllt 243

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